

IgE Fc C ϵ 3-C ϵ 4
Closed form



Fig. 1a

IgE Fc C ϵ 3-C ϵ 4
Open form

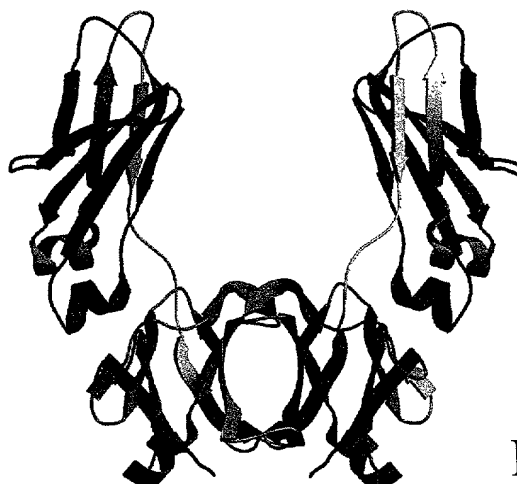


Fig. 1b

IgG Fc

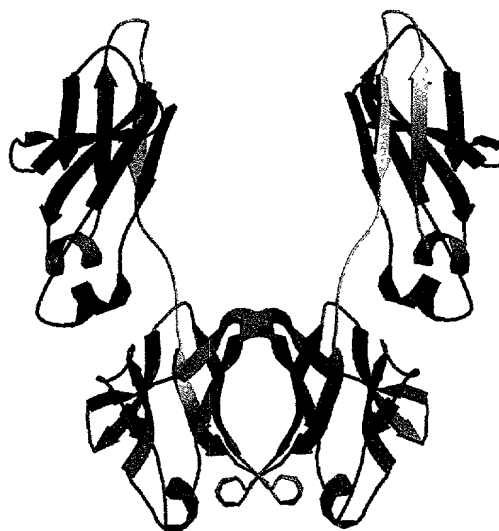


Fig. 1c

Closed
IgE-Fc

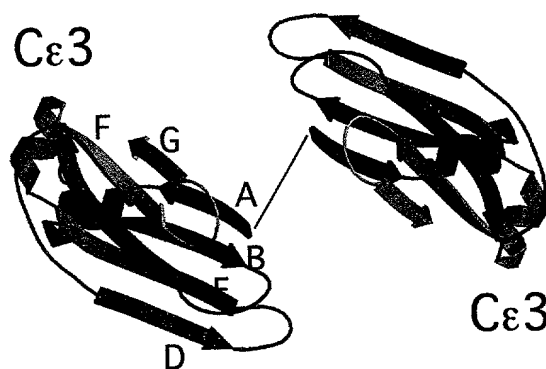


Fig. 2a

Open
IgE-Fc

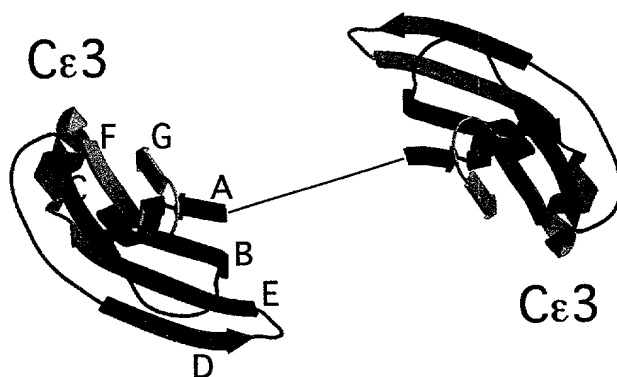


Fig. 2b

IgG-Fc

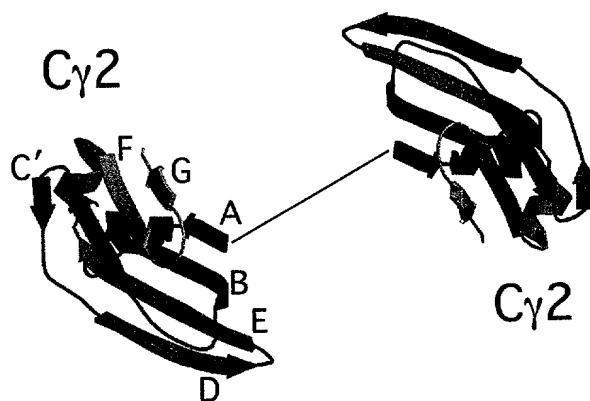


Fig. 2c

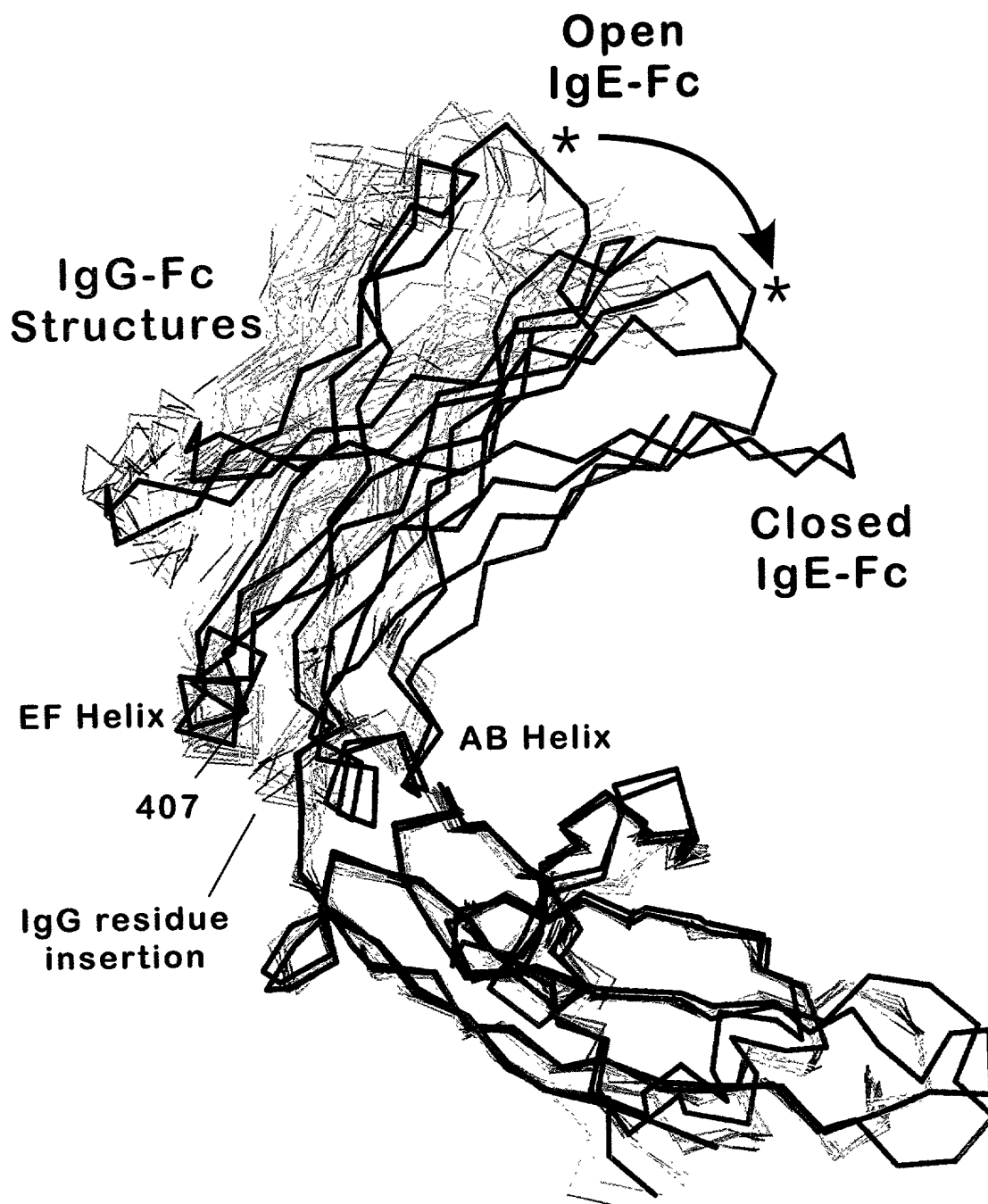


Fig. 3

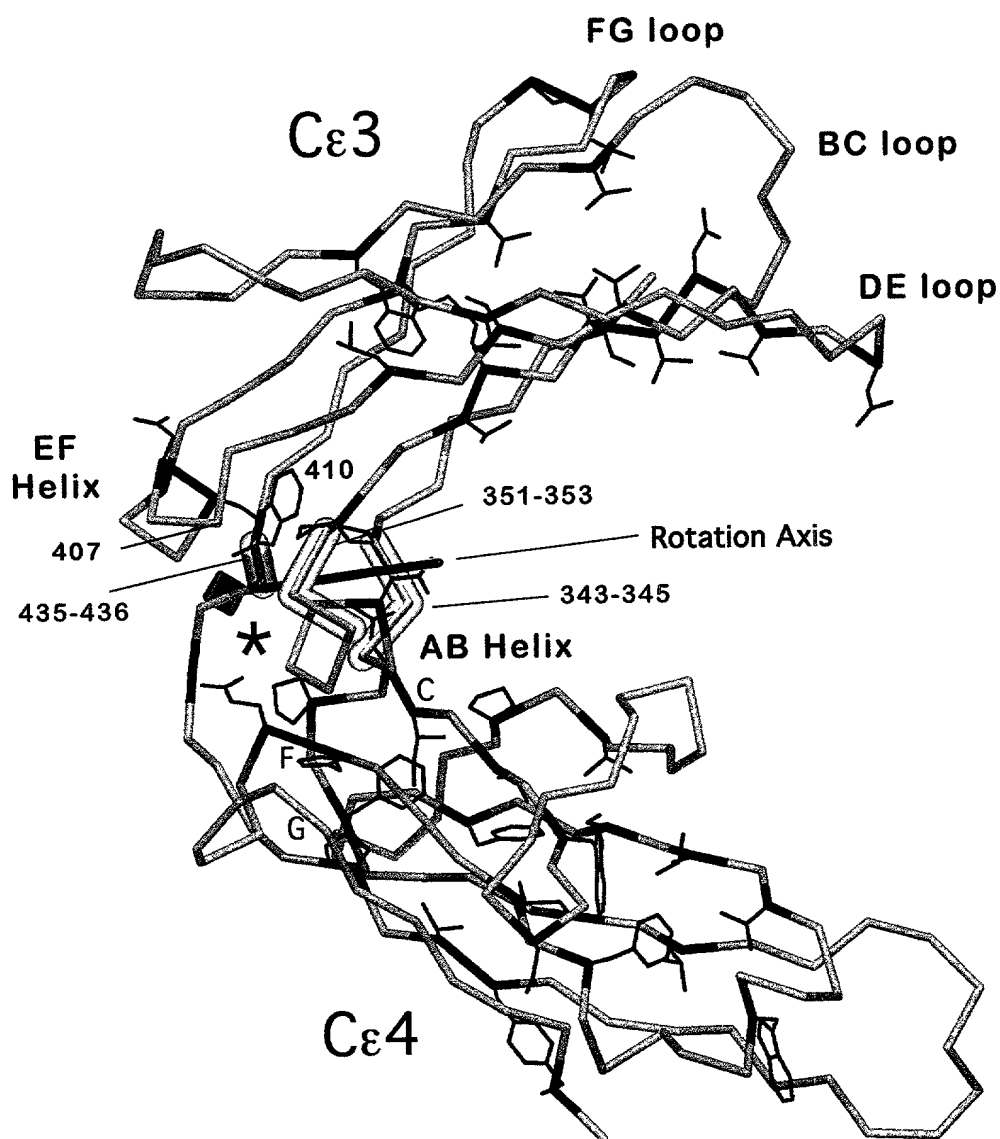


Fig. 4a

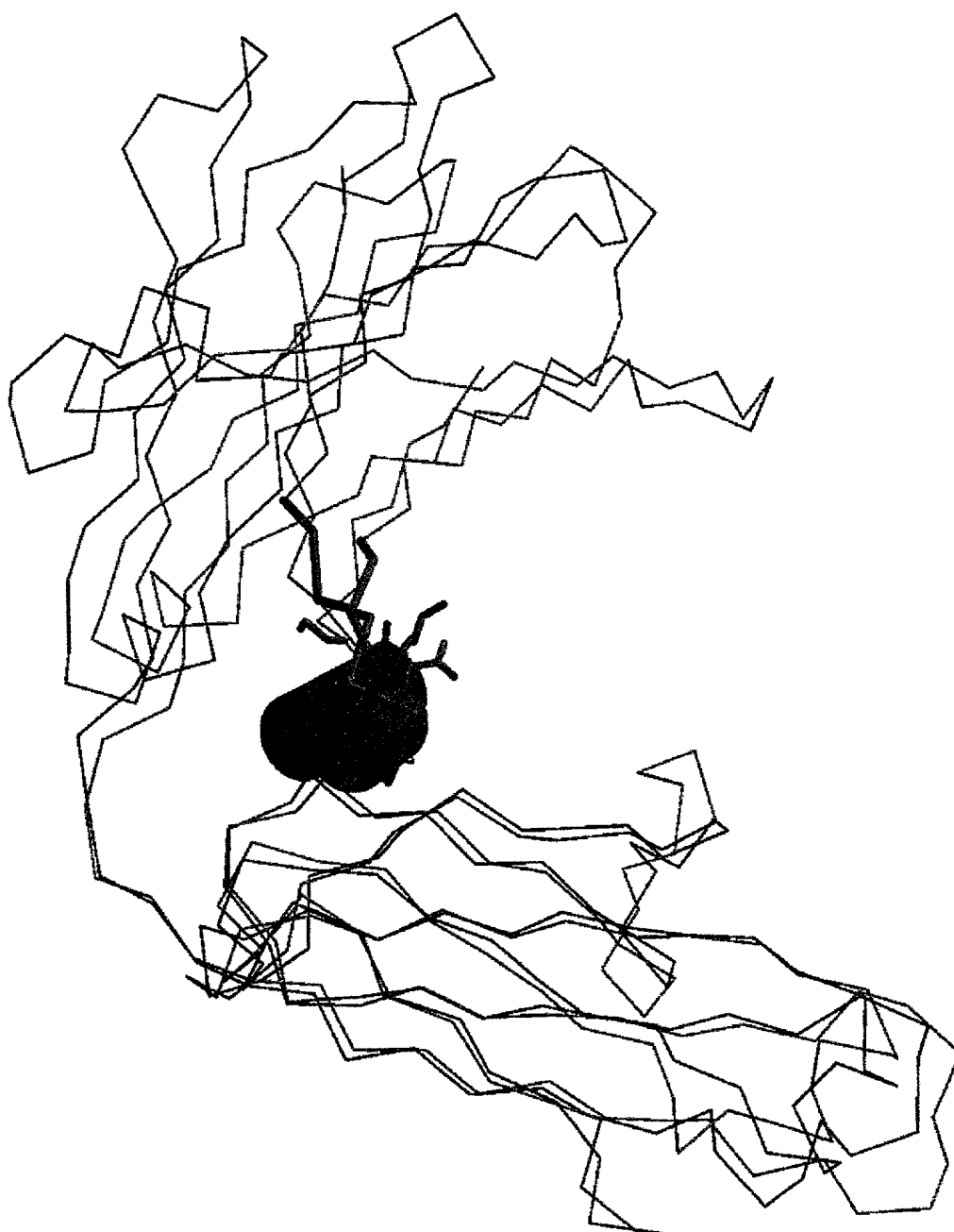


Fig. 4b

IgE-Fc $C\alpha$ - $C\alpha$ Distances (open-closed)

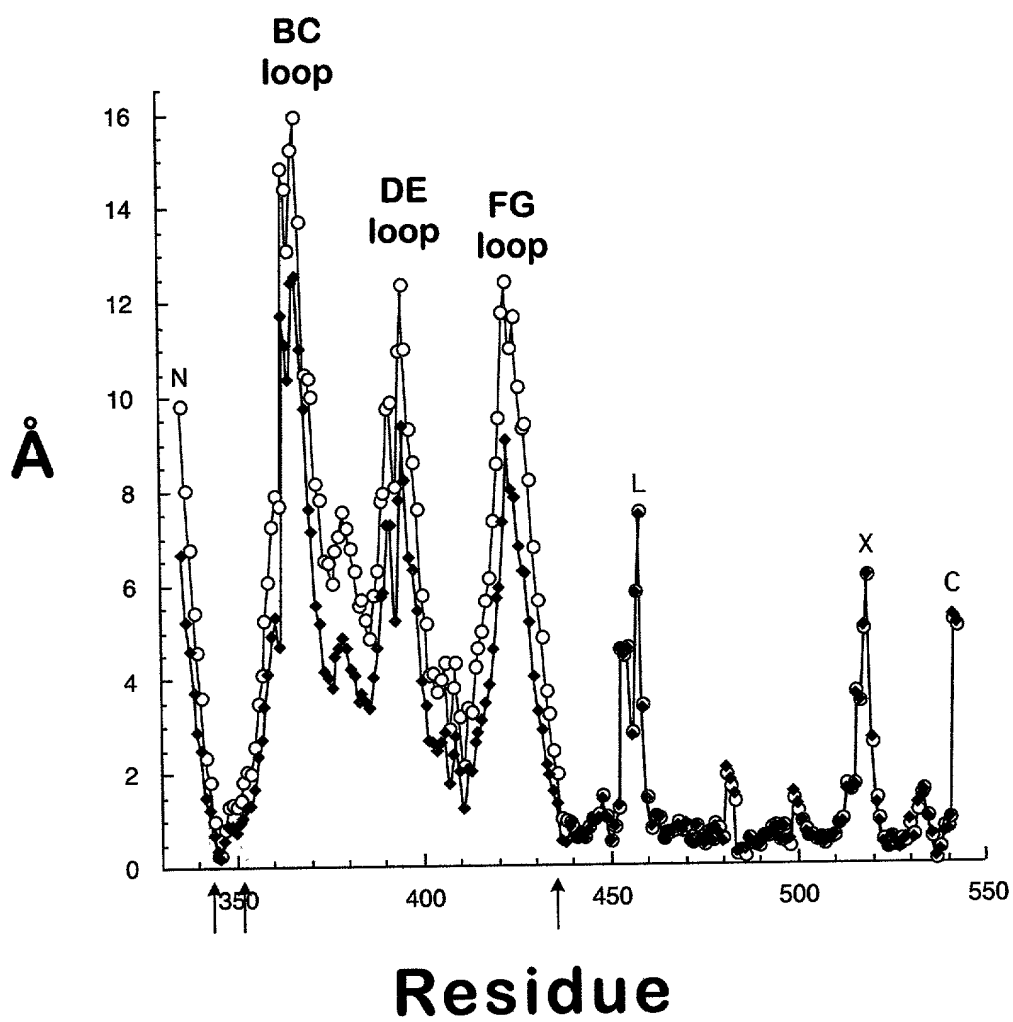


Fig. 4c

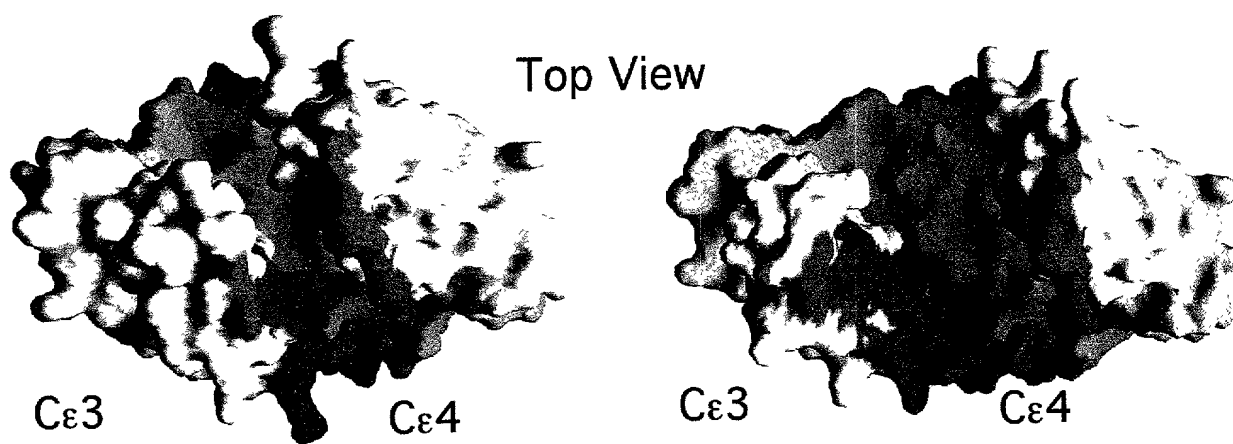


Fig. 5a

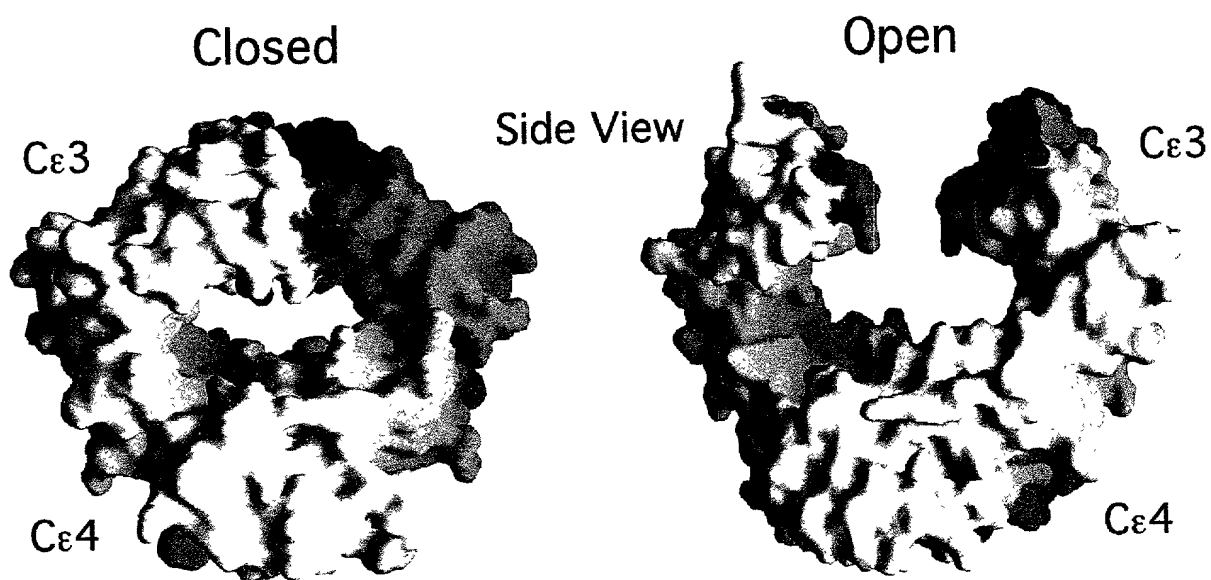


Fig. 5b

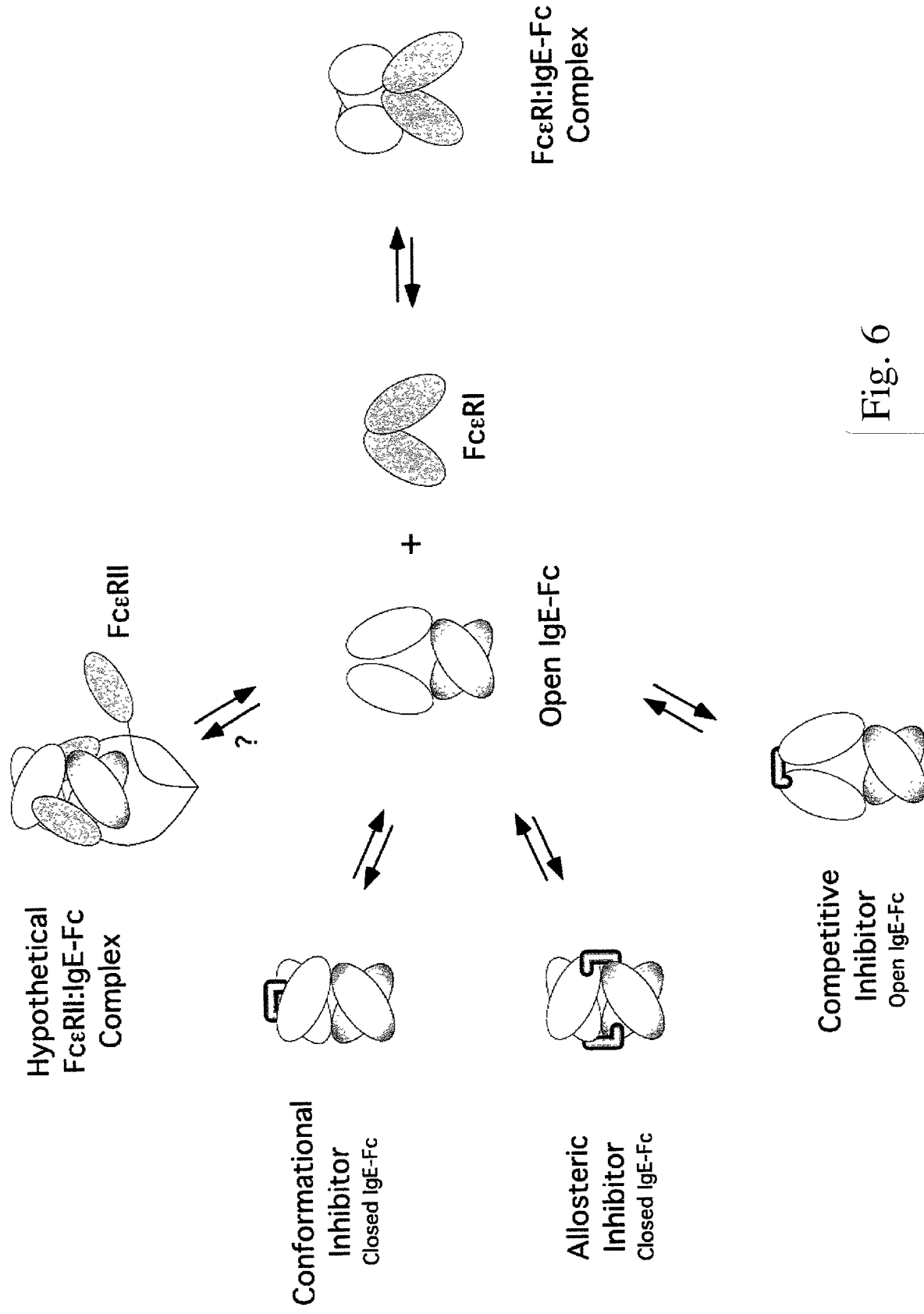
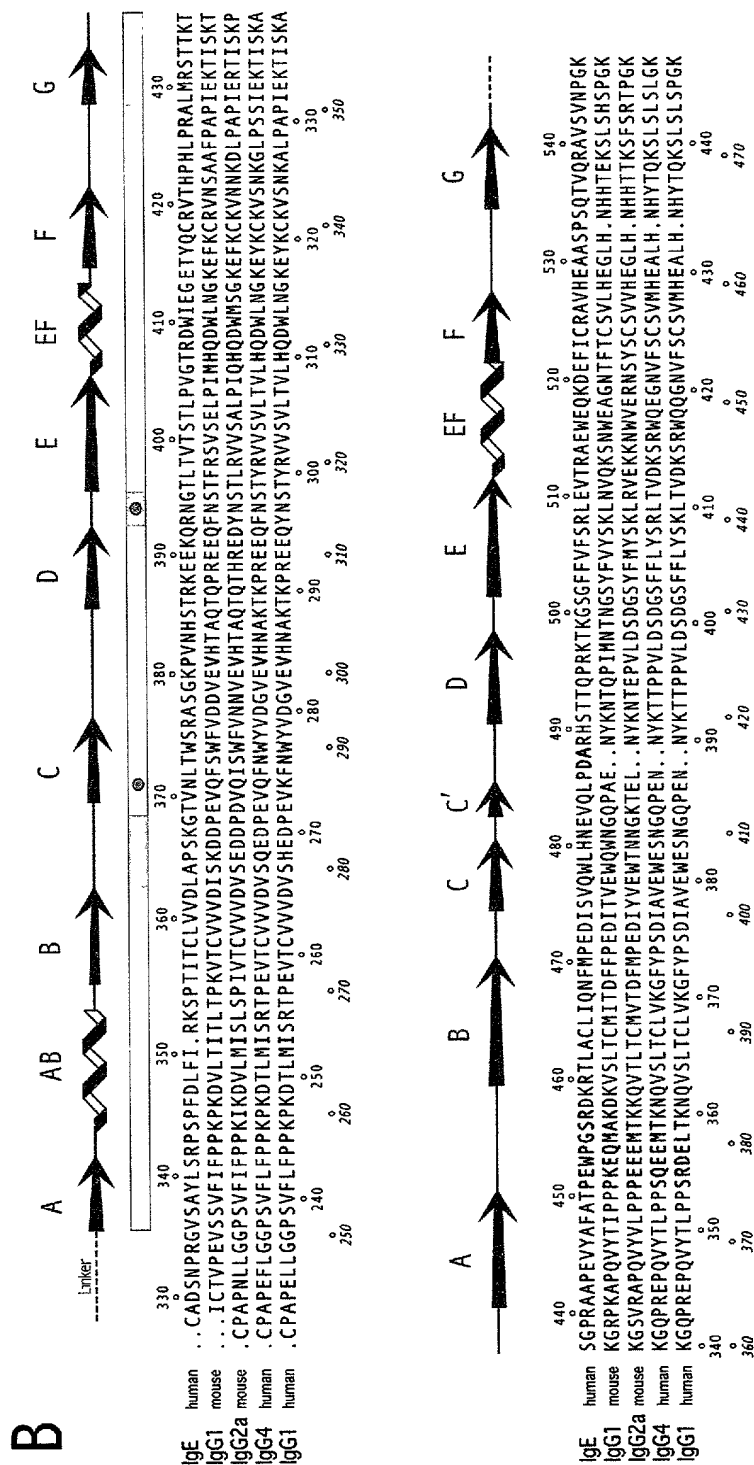
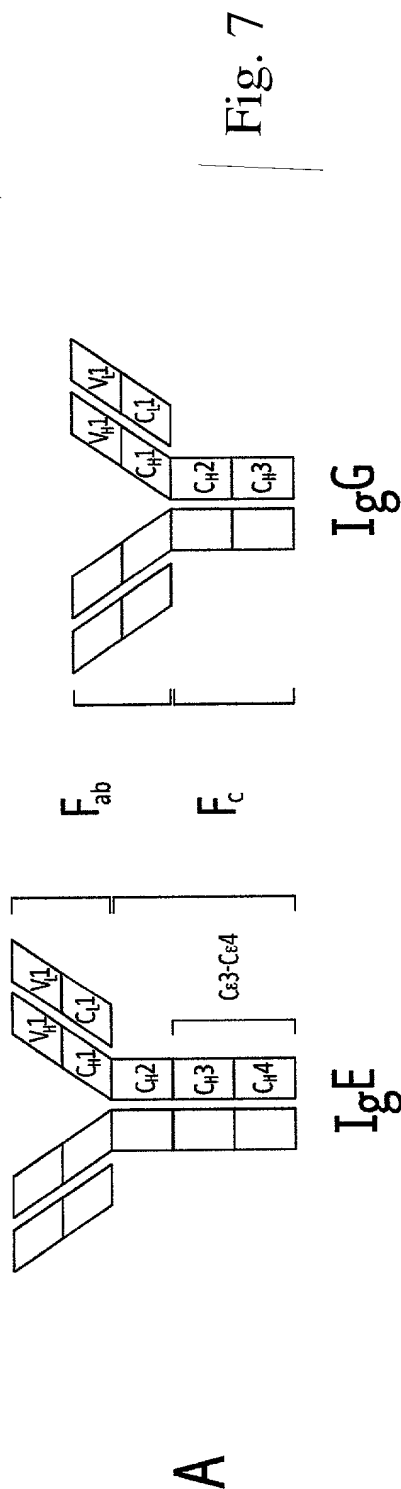


Fig. 6



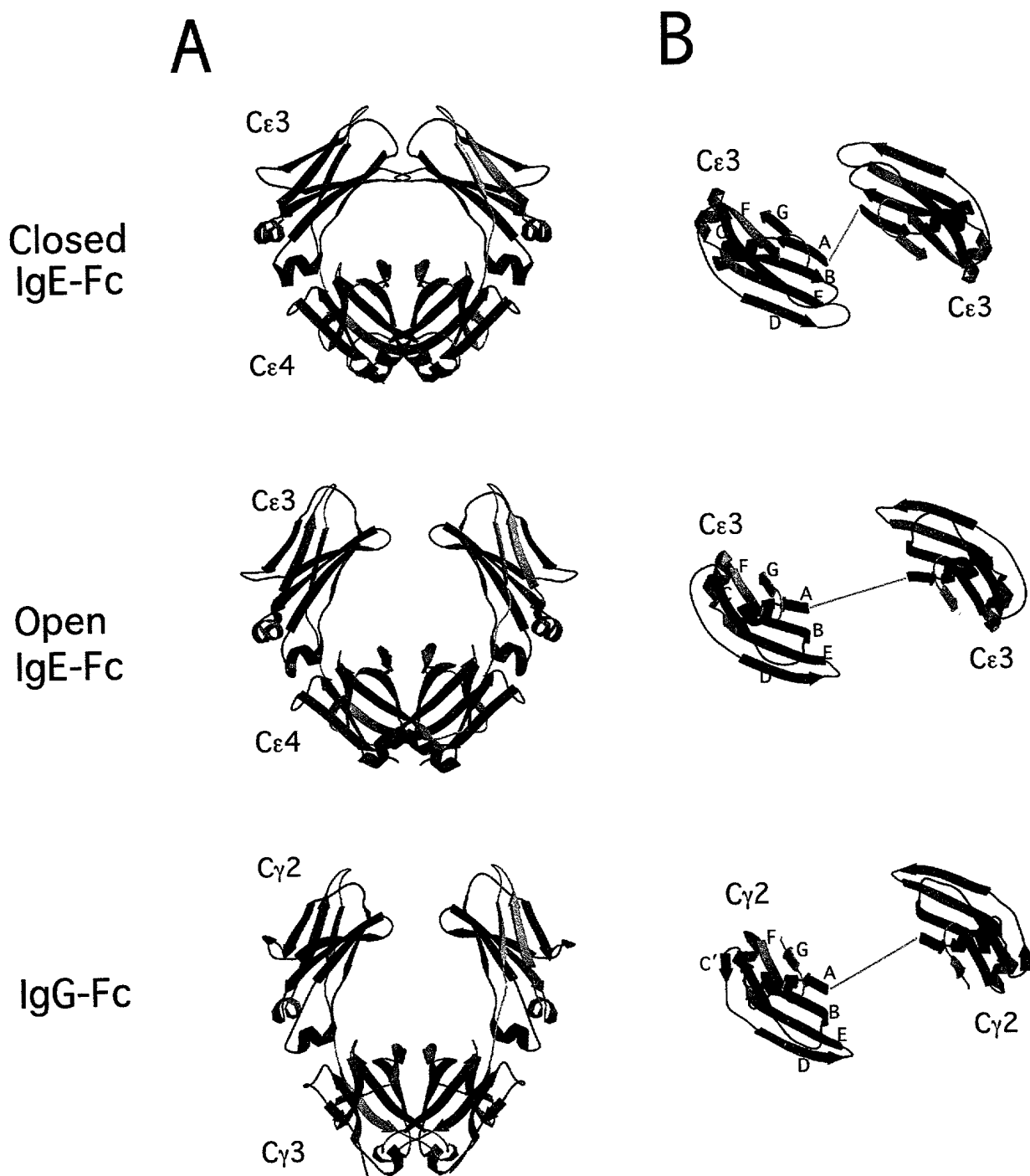


Fig. 8

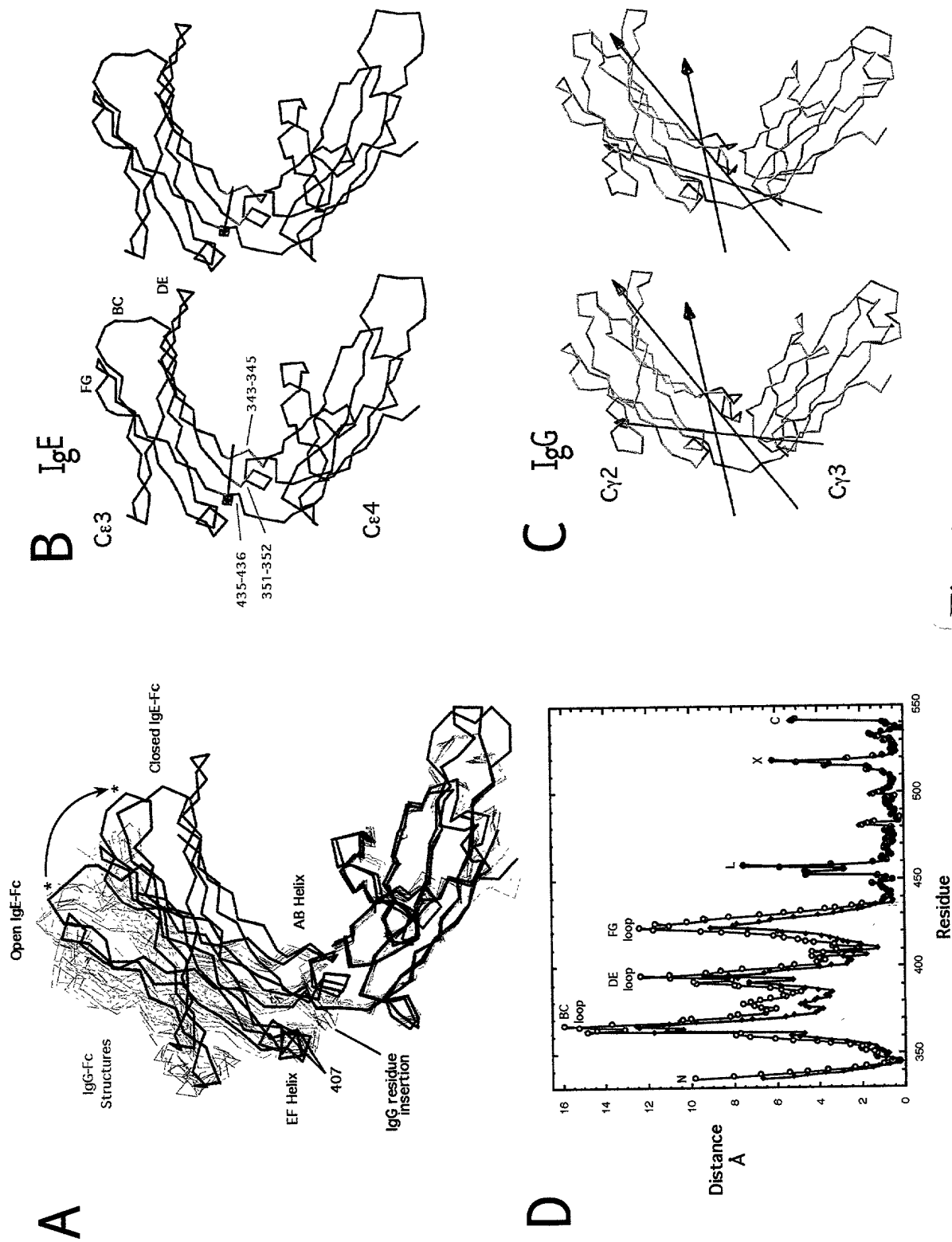


Fig. 9

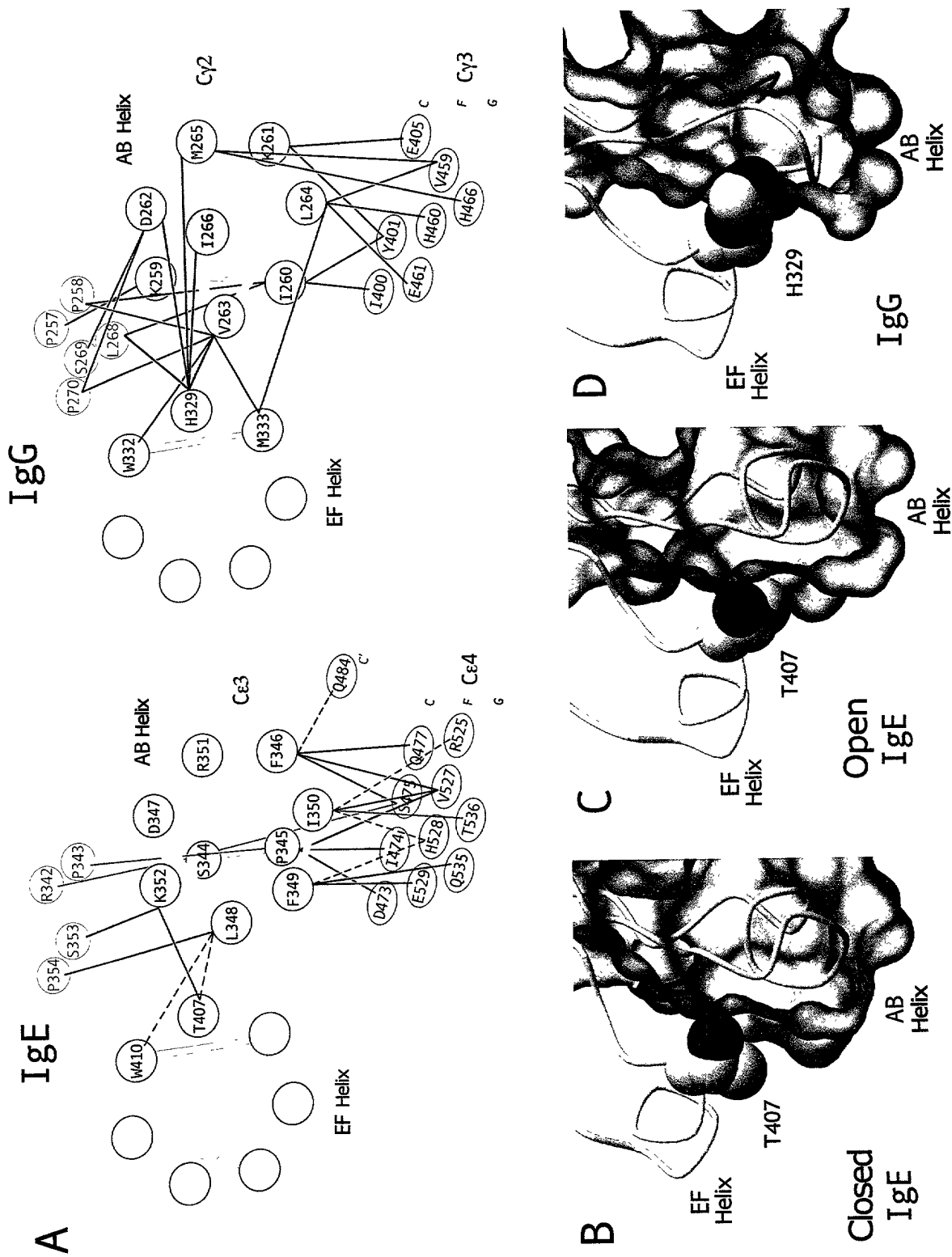


Fig. 10

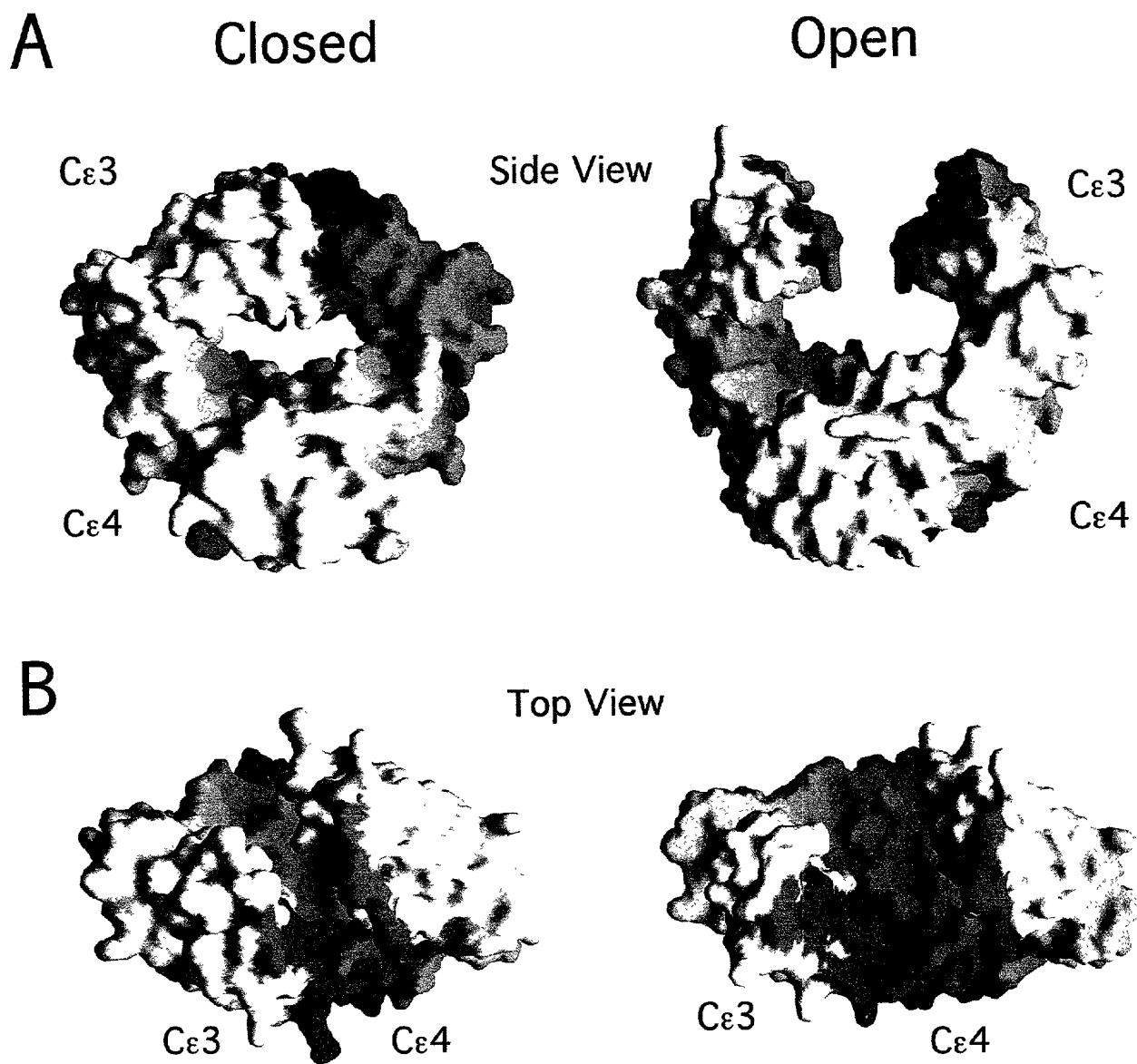


Fig. 11

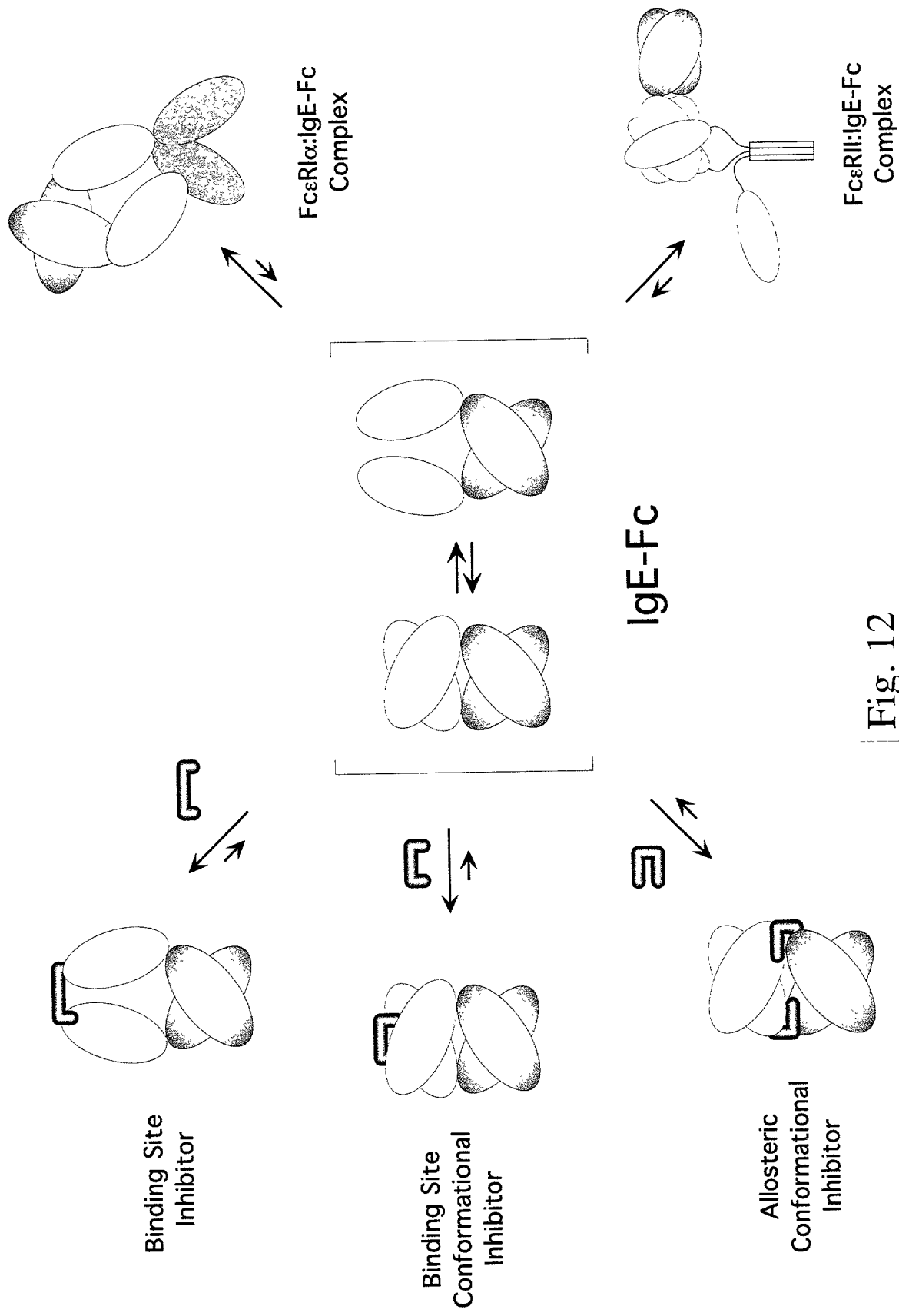
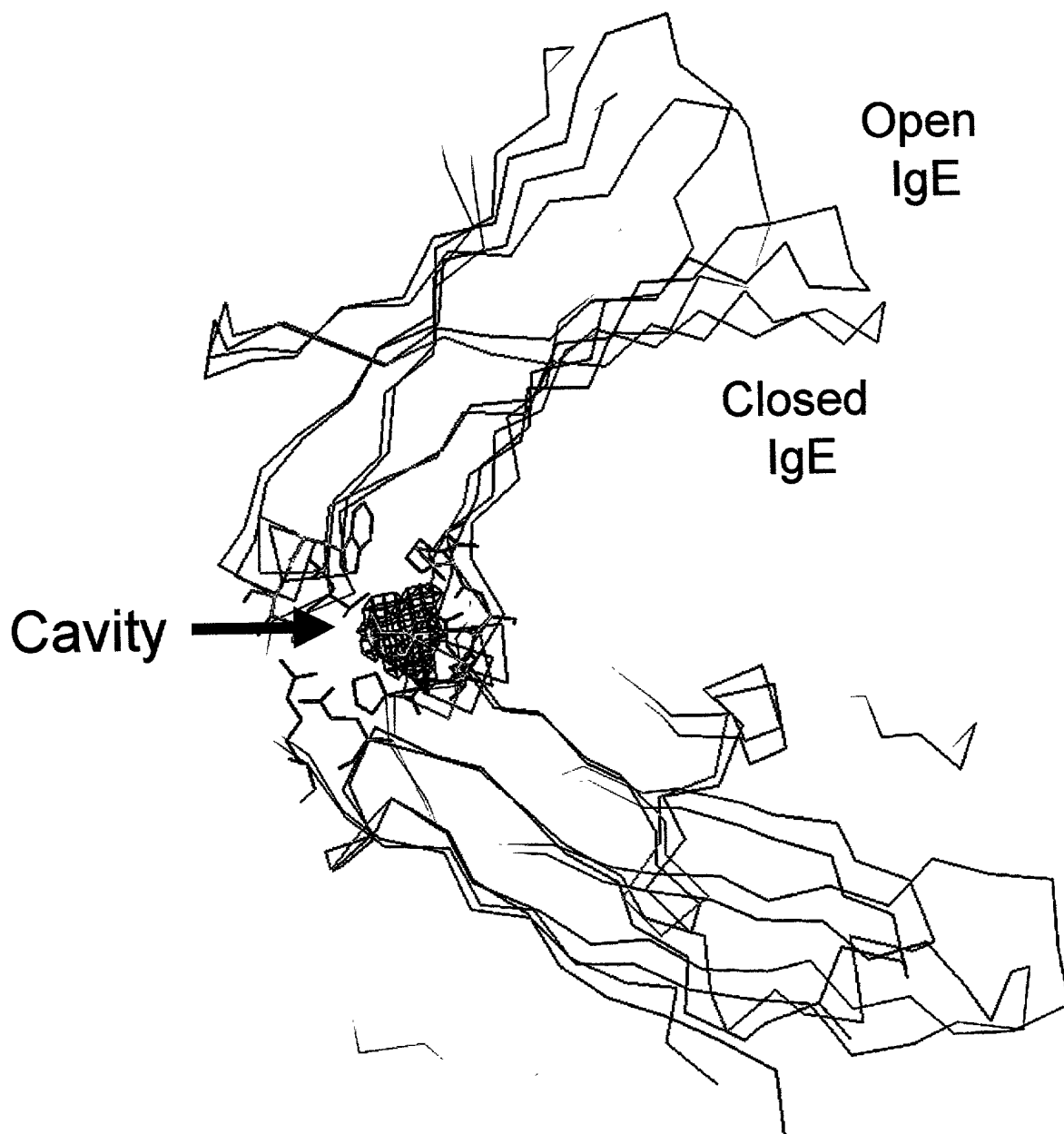


Fig. 12

A Potential Drug Binding Site Near the IgE-Fc Hinge



Surrounding residues include:
R342, P343, S344, P345, L348, W410, I411,
K435, T436, R440, P471, E472, D473, E529

Fig. 13